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Vol. 149, No. 3, 1987
December 31, 1987BIOCHEMICAL AND BIOPHYSICAL RESEARCH COMMUNICATIONS
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EXPRESSION OF CHIMERIC RECEPTOR COMPOSED OF IMMUNOGLOBULIN-DERIVED V. REGIONS AND T-CELL RECEPTOR-DERIVED C. REGIONS

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Received November 12, 1987

SUMMARY: Chimeric genes composed of immunoglobulin (Ig)-derived variable (V) regions and T-cell receptor (TCR)-derived constant (C) regions were constructed. The V_L and V_H genes showing anti-phosphorylcholine (PC) activity were used in this study. Two pairs of chimeric genes, V_L-C_β and V_H-C_α genes, and V_L-C_α and V_H-C_β genes, were inserted into an expression vector containing both ECO ϕ X174 and neo genes, and transfected into EL4 cells. Cells which express both chimeric receptor molecules were established. The activity of the transformants to the antigen was examined by using stopped-flow fluorimetry. An increase in the concentration of cytoplasmic calcium ion was observed after addition of Staphylococcus pneumoniae R36A bacteria grown in the choline-containing medium which express PC molecules, but not after the PC-negative bacteria grown in the ethanolamine-containing medium.

TCR is a heterodimer of disulfide-linked α and β chains (1). The essential features of both chains are rather similar to those of Ig(2). N-terminal regions of both chains are variable and form a dimeric antigen-combining site (3). In contrast to the recognition of antigen by antibody, where the antibody can bind to free antigen, the TCR does not recognize antigen alone but only in association with MHC molecule (4). To determine whether the difference in antigen recognition between T and B

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Abbreviations: Ig, Immunoglobulin; TCR, T-cell receptor; V, Variable; C, constant; H, heavy; L, light; PC, phosphorylcholine; ECO-MOL, ecotropic Moloney virus; LTR, long terminal repeat.

0006-291X/87 \$1.50

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cells derives from the structural difference in V regions of TCR and Ig, we constructed chimeric genes composed of Ig-derived V regions and TCR-derived C regions. The V_L and V_H gene showing anti-PC activity were used in this study. The V genes of the light (L) and heavy (H) chains of anti-PC myelomas and hybridomas have been well characterized (5,6). Only one germline V_H gene (5) and three germline V_L genes (6) are responsible for the anti-PC specificity. Cells which express chimeric receptor molecules were established. It is known that the activation of T lymphocytes is accompanied by an increase in the concentration of cytosolic calcium (7). The activity of the transformants to the antigen was examined by using stopped-flow fluorometry (8). The chimeric receptor molecules expressed on the transformants had the capacity to react with PC antigen and trigger T cell activation.

MATERIALS AND METHODS

Active V_H gene was isolated from myeloma TEPC15 by using J_H gene-containing DNA fragment as a probe (5). Active V_L gene isolated from myeloma S107 was donated by M.D. Scharff (Albert Einstein College) (9). The C genes of T-cell receptor α and β chains were isolated from C3H and C57BL/6 mouse DNA by using cDNAs of α and β chains (2,10) donated by M. Davis (Stanford University) as probes, respectively. The ecotropic Moloney virus (ECO-MOL) long terminal repeat (LTR) fragment was donated by M. Ishimoto (Kyoto University) (11). Plasmid PAGE145 was described previously (12). The bacterin and the eukaryotic cells containing this plasmid show kanamycin resistance at a concentration of 10 μ g/ml and G418 resistance at a concentration of 1 mg/ml, respectively. The nucleotide sequence of the constructed plasmids was determined by the M13-dideoxy method (13). EL4 cell was obtained from T. Taniguchi (Osaka University). Transfection experiments were done by protoplast fusion as described (14). Total RNA was extracted from cells by the Cuanidinium-CsCl method (15). Poly A-containing RNA was purified with the oligo (dT) cellulose column (15). Northern hybridization was performed as described (16). Anti- V_H antibody (17) and anti-T15 idiotype antibody (18) were donated by D. Civol (Israel) and C. Heusser (Basel), respectively. Stopped-flow fluorometry was carried out as follows. Four 2-loaded T-cells (19) (final 10⁶ cells/ml) were mixed with the target bacteria (final 4 \times 10⁸ cells/ml) at 37°C. Stopped-flow fluorescence was measured with a Union Giken stopped-flow spectro-photometer RA401 in combination with a microcomputer RA-450 system (8). Excitation wavelength: 335nm. We used a Hoya Y46 cut filter (which allows the emitted light with wavelength longer than 460 nm to come into the detector) in the fluorescence measurements.

RESULTS

Figure 1 shows the construction procedures for four kinds of chimeric genes: V_L - C_α , V_H - C_α , V_L - C_β and V_H - C_β . To express the chimeric genes in T cells, we used the ECO-MOL LTR as a transcriptional promoter. Since translation begins at the 5'-proximal AUG triplet in eukaryotic mRNAs (20), we determined the nucleotide sequence between the promoter and

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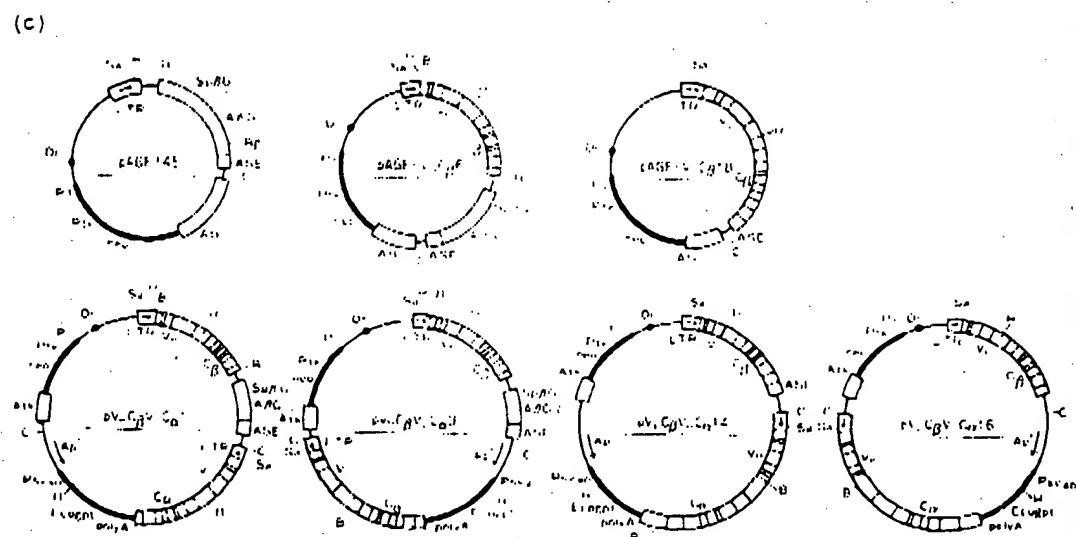
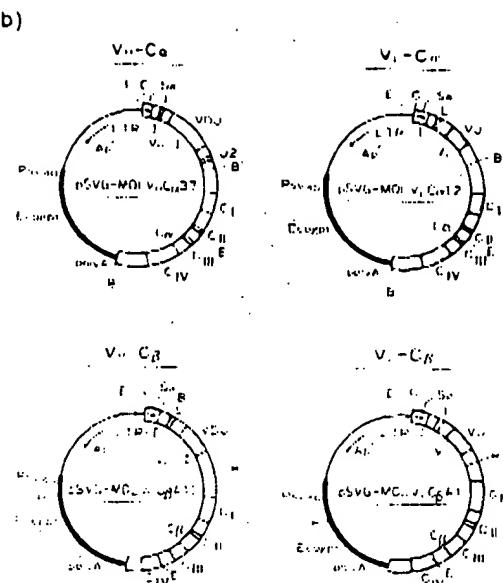
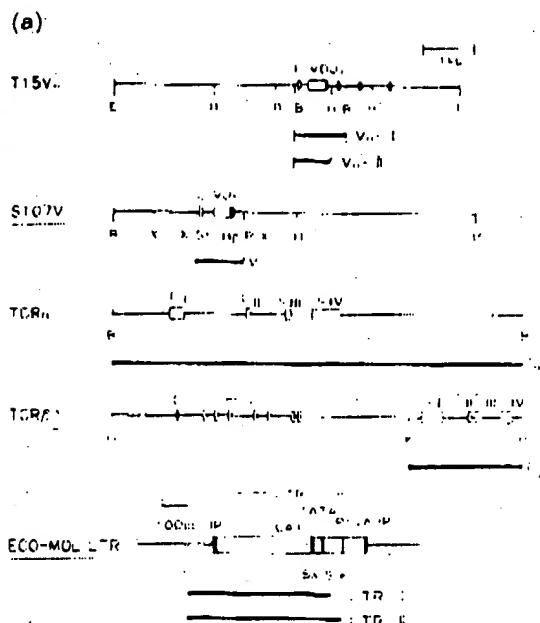


Fig. 1. Construction of the chimeric genes. (a) Restriction maps of the clones. The DNA fragments used for the construction of plasmids are indicated by thick lines. Restriction Enzymes: E, EcoRI; B, BamHI; H, HindIII; X, XbaI; S, SstI; Sp, SphI; P, PstI; K, KpnI; C, C1aI; S, SacI; s, SmaI. (b) For the construction of pSVG-MOLV C_H37, the C1aI-SmaI fragment of the ECO-MOLLTR(LTR-I) was connected to the BamHI-BamHI fragment of T15V_H (V_H-I), after the BamHI site at the 5' upstream of V_H gene was changed to a SphI site with Klenow enzyme. The resulting C1aI-BamHI fragment was connected to the BamHI-BamHI fragment of TCR_H chain as in the same polarity, and inserted into plasmid pSV2gpt. The EcoRI-C1aI region at the upstream of the LTR sequence was derived from pBR322 DNA. For the construction of pSVG-MOLV C_H41, the LTR-I fragment was connected to the SstI-BglII fragment of S107V_H (V_H). The resulting fragment was connected to the BamHI fragment of C₀ gene after changing the BglII site into BamHI site by using a BamHI linker, and inserted into plasmid pSV2gpt. For the construction of pSVG-MOLV C_H41, the C1aI-Kpn fragment of the ECO-MOL

the leader of ATG sequences for transformants cell(21), we used MOLV C_H (neo), protoplast f medium (23). gpt and neo pV_H C₀ V_L C_m chimeric genes in the gpt and transfected respectively. transformants Northern hybridization cells express EL4-5 contains neo resistant 1.45 kb with t

LTR(LTR-II) after changing enzyme, where fragment was changing respectively. the construct pSVG-MOLV C_H instead of whose structure resistance to pSV2gpt. MOLV C_H (neo) construct DNA, PAGE14% polylinker designation splicing site thymidine kinase of PI and MOLV C_H All site of PAGE pSVG-MOLV C_H resulting BglII fragm pSVG-MOLV C_H PAGE-V_L C_m C1aI-digeste pV_L C_H C_m pV_L C_H C_m The overall hybridizatio sizes of each

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the leader of V_H and V_L genes in the constructed plasmids. There were no ATG sequences in the 5' upstream regions (Fig.2). In an initial trial for transformation experiments, the EL4 cells, presumed to be a helper T cell (21), were transfected with two plasmid DNAs: pSVC-MOLV V_H C_3 and pSVC-MOLV V_L C_B (neo), or pSVC-MOLV V_L C_12 and pSVC-MOLV V_H C_B (neo), simultaneously, by protoplast fusion (14) and selected in the gpt (22) and neo selection medium (21). However, we did not obtain any transformants showing both gpt and neo resistance. We constructed the plasmids: pV H_B V L_B 1, pV H_B V L_B 3, pV L_B V H_A 14 and pV L_B V H_A 16, containing two chimeric genes in each plasmid DNA (Fig.1c). From 4×10^6 EL4 cells, in the gpt selection medium, we obtained 7, 22, 13 and 6 transformants transfected with pV H_B V L_B 1, pV H_B V L_B 3, pV L_B V H_A 14 and pV L_B V H_A 16, respectively. In the neo selection medium, we obtained 2, 10, 9 and 3 transformants transfected with the above plasmid DNAs, respectively. By Northern hybridization (16) with the V_H and V_L probes, we searched for cells expressing both chimeric genes, and obtained two transformants: EL4-5 containing pV H_B V L_B 3 and EL4-14 containing pV L_B V H_A 14, as neo resistant cells. EL4-5 gave a band at 1.35 kb with the V_H probe and at 1.45 kb with the V_L probe (Fig.3). EL4-14 gave a band at 1.45 kb with

1. LTR(LTR-II) was connected to the BamHI-HindIII fragment of T15V(V_H -II) after changing the KpnI and the BamHI sites into flush ends with Klenow enzyme, which creates a KpnII site at the junction point. The resulting fragment was connected to the KpnI-HindIII fragment of β ₂ gene after changing the KpnI and HindIII sites into HindIII and KpnII sites, respectively, by using linkers, and inserted into plasmid pSV2gprt. For the construction of pSVC-MOLV V_L C_12 , the protocol was the same as that of pSVC-MOLV V_H C_3 except for the change of the BglII site into HindIII site instead of KpnII site. Independently, we constructed four other plasmids whose structures are essentially the same as shown in (b) except for neo resistance gene in place of Ecogpt gene by using plasmid pSV2neo instead of pSV2gprt. We named them pSVC-MOLV V_H C_3 (neo), pSVC-MOLV V_L C_12 (neo) and pSVC-MOLV V_H C_B (neo), respectively. (c) For the construction of the plasmids containing two chimeric genes in one plasmid DNA, pAGE145(3) was used as a vector. The LTR is the ECO-MOL-LTR and a polylinker shown in Fig.2 was inserted into the SmaI site. The designations are as follows: SpRC, ARG, ASE and Ark are β -globin splicing signal, β -globin polyA signal, SV40 early gene polyA signal and thymidine kinase polyA signal, respectively. PI and Ptk are the promoters of PI and thymidine kinase genes. The BamHI-BamHI fragment of pSVC-MOLV V_H C_3 containing the chimeric V_H - C_3 gene was inserted into the BamHI site of pAGE145, resulting in the plasmid pAGE-V H C_3 . The Clal-digested pSVC-MOLV V_L C_12 was connected to the Clal-digested pAGE-V H C_3 , resulting in two clones: pV H_B V L_B 1 and pV H_B V L_B 3. The SacI-BglII fragment of pAGE145 was replaced by the SacI-BamHI fragment of pSVC-MOLV V_L C_12 containing the chimeric V_L - C_12 gene, resulting in plasmid pAGE-V L C_12 . The Clal-digested pSVC-MOLV V_H C_B was connected to the Clal-digested pAGE-V L C_12 , resulting in two plasmids: pV L_B V H_A 14 and pV L_B V H_A 16. The sizes of the four plasmids: pV H_B V L_B 1, pV H_B V L_B 3, pV L_B V H_A 14, and pV L_B V H_A 16 are 23, 23, 21.6 and 21.6 kb, respectively. The overall structures are confirmed by restriction mapping, Southern hybridization and nucleotide sequencing of the connected regions. The sizes of each plasmid and gene shown in (b)(c) and (d) are arbitrary.

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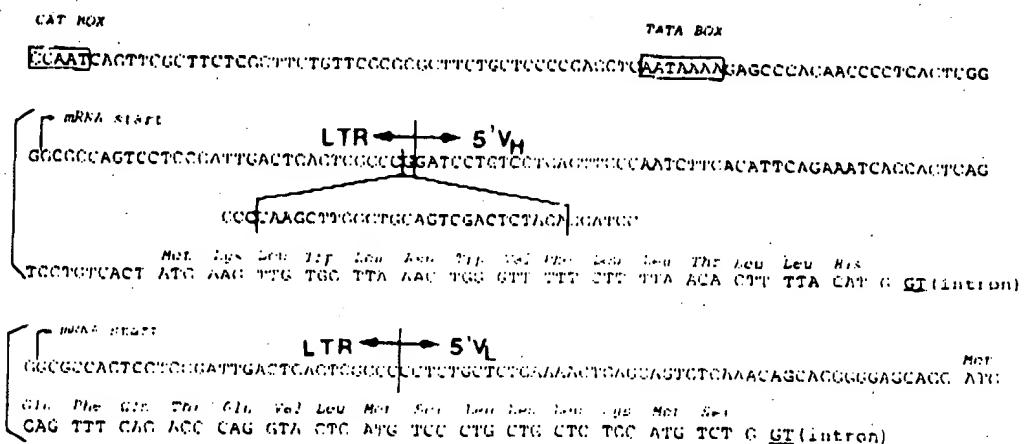


Fig.2. Nucleotide sequence from the promoter region to the first intron of the constructed V_H and V_L genes. The nucleotide sequence of the promoter was as described (1). The junction points of the LTR and V genes are at the Sma I or Kpn I site in the ECO-M01 LTR, the Bam H site in the 5' upstream of T15 V_H gene and the Stu I site in the 5' upstream of S107 V_L gene as described in the legend of Fig.1. Since a polylinker was inserted into the Sma I site in plasmid pAGE14S (12), there are 26 extra nucleotides at the boundary of the LTR and 5' V_H in p λ - V_H ; b. The determined sequences are the same as the published sequence (5) except for a two nucleotide deletion in the 5' untranslated region of the V_H gene. The position of the initiation codon in S107 V_L gene is estimated from the locations of the octamer ATTTGCAT (35) and TATA-like sequences whose positions were removed from the constructed plasmids.

the V_H probe and at 1.25 kb with the V_L probe. The size of each band indicates that RNA splicing occurred between an Ig-derived J gene and a TCR-derived C gene. The size differences should reflect those of the 3' end structure of the constructed genes (Fig.1).

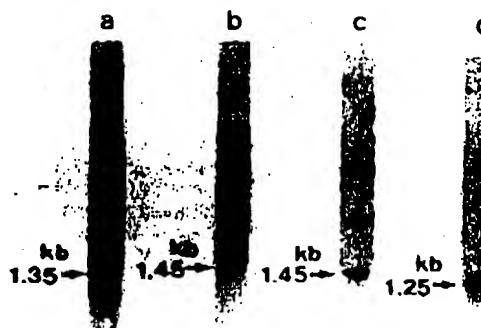


Fig.3. Northern hybridization of the transformants with the V_H and V_L probe. PolyA-containing RNA was prepared from each transformant. Two micrograms of RNA were subjected to 1% agarose gel electrophoresis. The size was estimated using 28S and 18S rRNA as size markers. Hybridization was carried out with (a)(c) the V_H probe (which corresponds to V_{HII} in Fig.1a), and (b)(d) the V_L probe (which corresponds to V_{L} in Fig.1a). (a)(b)EL4-5, (c)(d)EL4 14.

To determine chimeric receptor used two monoclonal idiotypic antibodies reacted with cross-reactivity (24), or some antibodies were a problem, we used antigen without

It is known that increase in the target cells, then, calcium influx. This increase is using stopped-source of PC as bacteria (25) molecules, and bacteria grown in both EL4-5 and the T cells will and that EL4 is bacteria. The presumably express PC antigen and t

The TCR is Lyt2/T8 and L3T4 interactions between groups (30,31). TCR alone may not allow the receptor allelic MHC protein difference in a computer analysis

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To determine whether translation of mRNAs and expression of the chimeric receptor molecules on the cell surface occurred in these cells, we used two monoclonal antibodies: anti- V_H antibody (17) and anti-T15 idiotype antibody (18). However, the EL4 used as a recipient cell itself reacted with anti-T15 idiotype antibody, presumably due to the crossreactivity to Thy-1 which shares the same epitope as T15 Idiotype (24), or some other surface molecules, and the results with anti- V_H antibody were ambiguous because of the high background. Because of these problems, we directly examined the activity of the transformants to the antigen without identification of the proteins.

It is known that the activation of T lymphocytes is accompanied by an increase in the concentration of cytosolic calcium (7). After binding to target cells, T lymphocytes first increase their membrane fluidity and, then, calcium is released from intracellular stores (8). After that, calcium influx occurs from the external medium into T lymphocytes (8). This increase in the concentration of cytoplasmic Ca^{2+} ion can be traced by using stopped-flow fluorometry (8) with a fluorescent probe. As the source of PC antigen, we used heat-killed Staphylococcus pneumoniae R36A bacteria (25) grown in the chorine-containing medium which express PC molecules, and as the negative control, we used heat-killed PC-negative bacteria grown in ethanolamine-containing medium (27). Figure 4 shows that in both EL4-5 and EL4-14 cells, calcium influx was observed after mixing the T cells with PC-positive bacteria but not with PC-negative bacteria, and that EL4 itself did not react with either PC-positive or PC-negative bacteria. These results indicate that the chimeric receptor molecules presumably expressed in the transformants have the capacity to react with PC antigen and trigger T cell activation.

DISCUSSION

The TCR is associated with the T3 complex (28). Furthermore, the Lyt2/T8 and L3T4/T4 surface glycoproteins are thought to be involved in the interactions between T cells and their target cells (29). However, two groups (30,31) have presented direct evidence that the α and β chains of TCR alone may define the dual, namely antigen and MHC, specificity. Transfection and expression of the genes for a particular α and β pair allow the recipient cell to respond to the combination of antigen and allelic MHC product recognized by the donor T cell (30,31). The difference in antigen recognition between B and T cells may originate in the structural difference in Ig V regions and TCR V regions, although computer analyses (32), based on published sequence data, indicate that the

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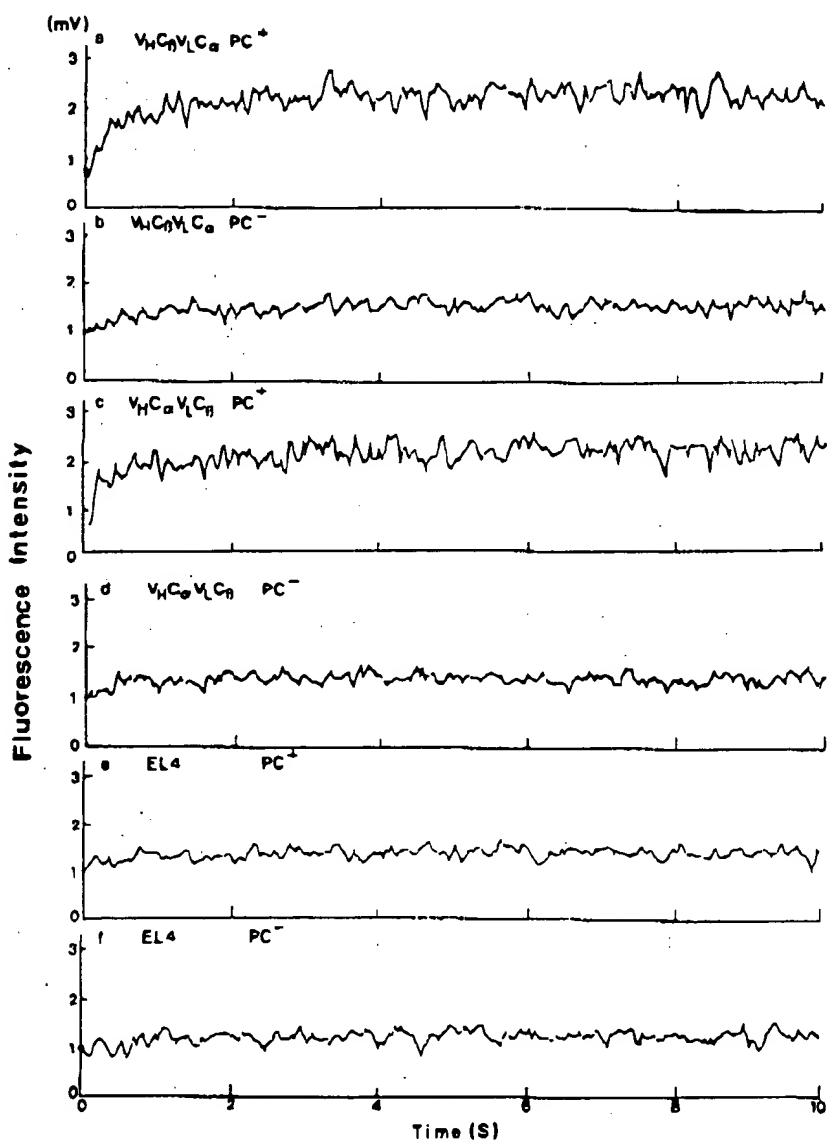


Fig. 4. Stopped-flow fluorometry traces indicating the effects of the antigens on the T-cells with chimeric receptors. The data are shown as the average of 15 times measurements.
 (a) Effect of PC^+ on the fluorescence of fura 2-loaded EL4-5.
 (b) Effect of PC^- on the fluorescence of fura 2-loaded EL4-5.
 (c) Effect of PC^+ on the fluorescence of fura 2-loaded EL4-14.
 (d) Effect of PC^- on the fluorescence of fura 2-loaded EL4-14.
 (e) Effect of PC^+ on the fluorescence of fura 2-loaded EL4.
 (f) Effect of PC^- on the fluorescence of fura 2-loaded EL4.

α and β chains of TCR are organized into immunoglobulin-like domains consisting of multistranded antiparallel β -sheet bilayers. Our present experiments indicate that the chimeric receptor composed of Ig-derived V regions and TCR-derived C regions has the capability to trigger T cell

activation with influx is an helper and cytotoxic ability of cells. In any antigens described in the

Recently, chimeric genes Although a chimeric gene can form an apparently secreted, it reason for for and V_L-C_β , a Ig's character without C domain.

We thank C. Heusser and I. We thank Dr. R. K. are grateful to transfection method for their encouragement the Ministry of Health Universit the Uehara Science

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activation with the antigen alone, without MHC molecules. Since calcium influx is an early transmembrane event (8), we are now examining whether helper and cytolytic functions occur with antigens. If this is the case, the ability of Ig to bind to free antigens could be transferred into T cells. In future, it might become possible for T cells recognizing any antigens without MHC restriction to be produced by the technique described in this paper.

Recently, Gascoigne *et al.* (33) reported the expression of reverse chimeric genes composed of TCR-derived V regions and Ig-derived C regions. Although a chimeric protein $V_{\alpha}-C_{Y2b}$ combined with a normal λ light chain to form an apparently normal tetrameric immunoglobulin molecule that was secreted, it did not combine with a chimeric $V_{\beta}-C_{\kappa}$ chain (33). The reason for formation of our chimeric chains: $V_{\alpha}-C_{\beta}$ and $V_{\beta}-C_{\alpha}$, and $V_{\alpha}-C_{\alpha}$ and $V_{\beta}-C_{\beta}$, although not directly demonstrated, might be related to the Ig's character in that V_{α} and V_{β} domains can be intrinsically associated without C domains to form an F_v fragment (34).

ACKNOWLEDGEMENTS

We thank Drs. M. Davis, A. Ishimoto, H. Seto, M.D. Scharff, D. Civoi, C. Heusser and M. Potter for providing us the materials used in this study. We thank Dr. R. Kubo for helpful discussions to prepare this manuscript. We are grateful to Drs. M. Matsukawa and T. Taniguchi for teaching us the transfection method. We also thank Drs. Y. Takagi, I. Ishiguro and K. Fujita for their encouragement. This work was supported in part by grants from the Ministry of Education, Science and Culture of Japan; Fujita-Gakuen Health University; Japan Private School Promotion Foundation (to Y.K.) and the Uehara Science Foundation (to Y.K.).

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Received November

SUMMARY. Transcription of the lipopolysaccharide mRNA for the early multiple pathways mRNA for JB was not (Intron 6) regulation of the present evidence to the JB gene. By induced transcription of the gene in LPS-treated cells, the gene is regulated mechanistic pathways upon the cell type. © 1987 Academic Press.

INTRODUCTION.
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Abbreviations
platelet-derived